



THE UNIVERSITY OF TEXAS

**MDAnderson**  
**Cancer Center**

Making Cancer History®

# **Basket Trials Revisited: Faster, Better, Smarter Trials. Precision Medicine: Insights, Challenges and Perspectives in Academia**

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# Disclosures

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# Initiative for Molecular Profiling in Advanced Cancer Therapy (IMPACT)

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## Hypothesis, 2007

- Selection of therapy based on patients' tumor molecular analysis will improve clinical outcomes compared to the standard approach

## Methods

- Patients who exhausted standard treatment options or had incurable rare cancers were referred to our Phase I program for treatment.
- CLIA-certified tumor molecular testing in consecutive patients referred for treatment.
- Genes analyzed: 1-50, depending on time of testing
- Trials available against various targets
- Treatment: matched targeted therapy, if available; if unavailable, non-matched.
- Retrospective analysis, exploratory.

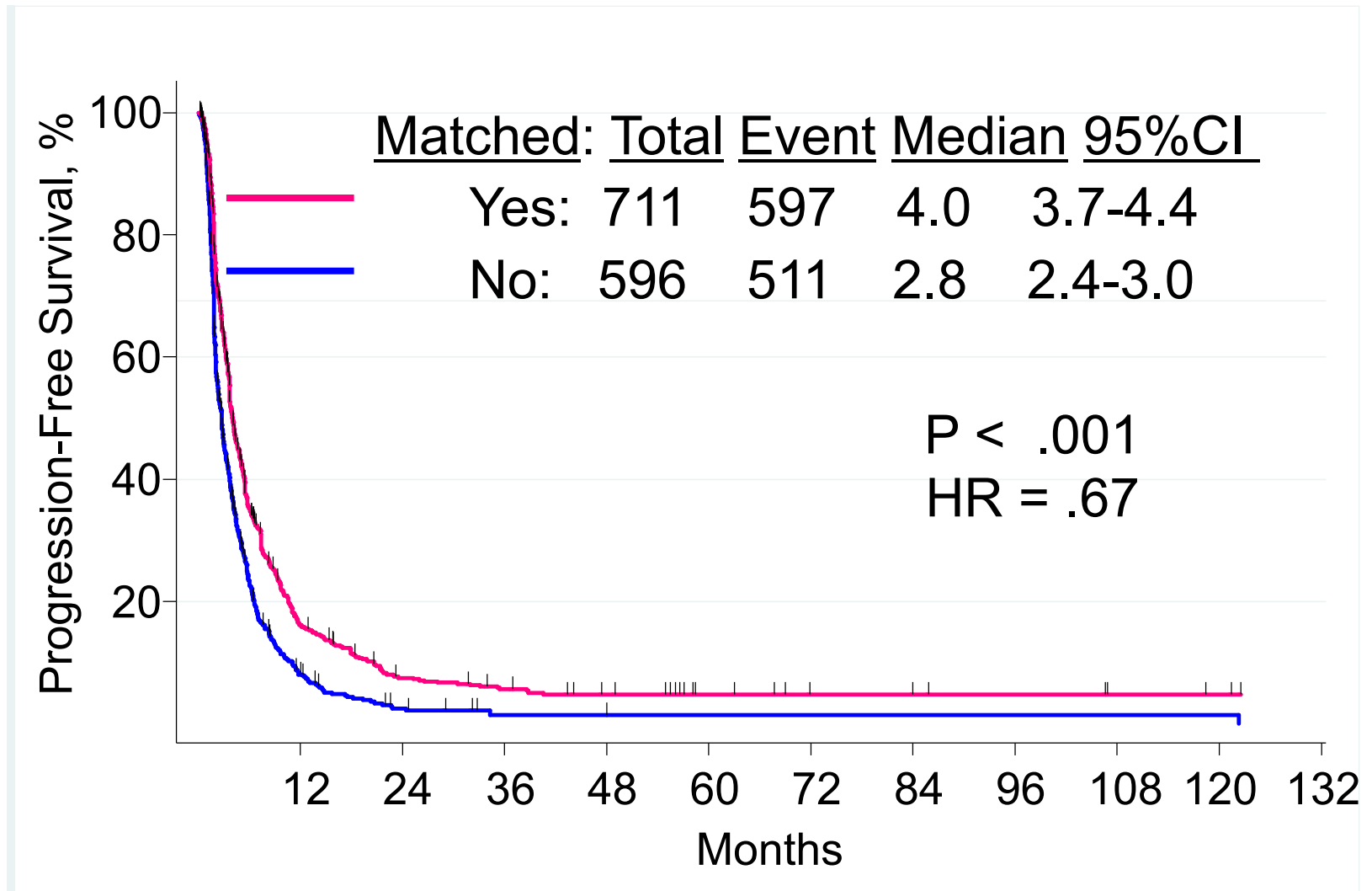
[www.clinicaltrials.gov](http://www.clinicaltrials.gov) NCT00851032

# IMPACT: Results

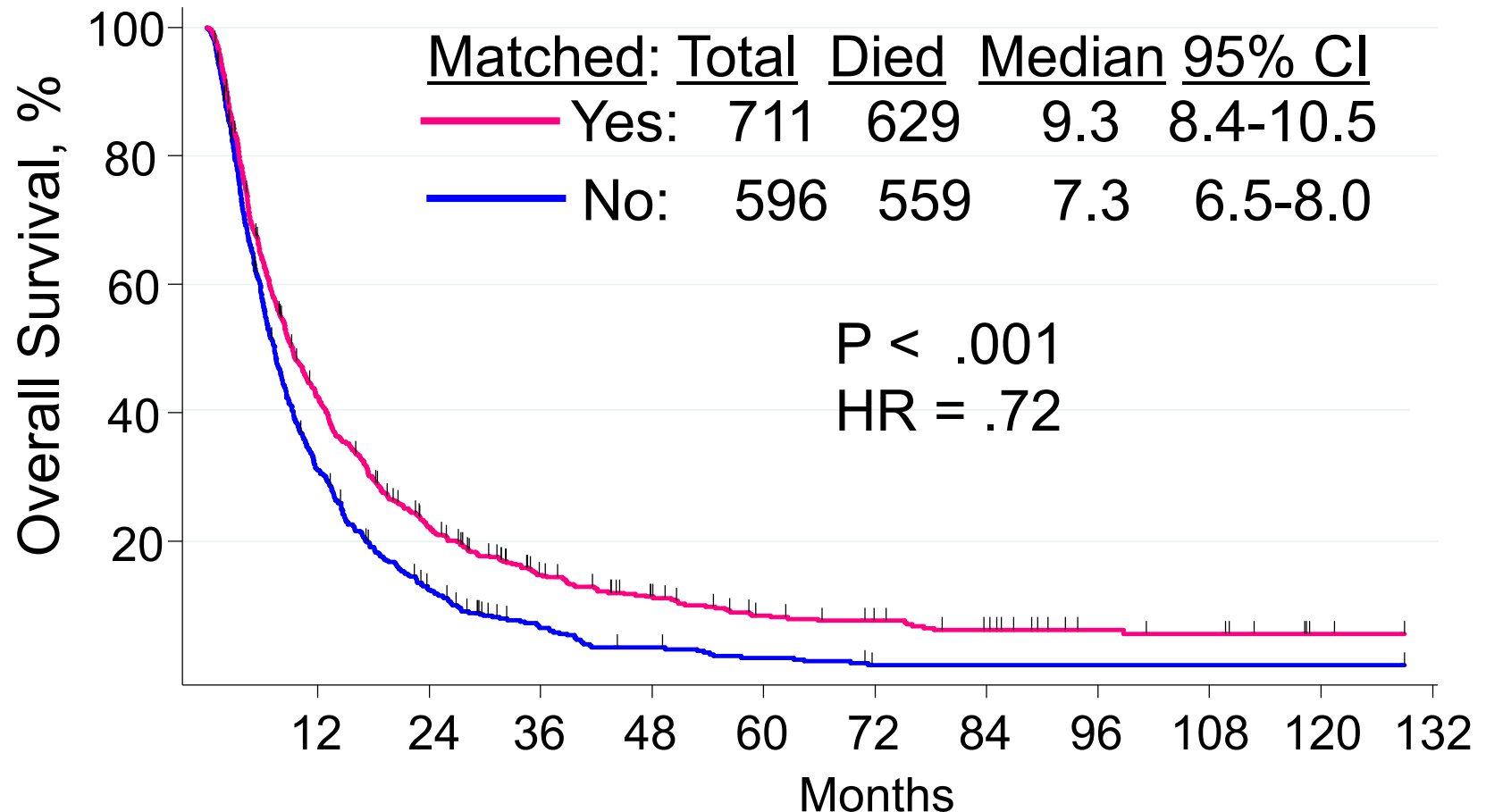
- Molecular testing: N = 3,743 (2007-2013)
- 1,307 (34.9%):  $\geq 1$  targetable molecular alteration
- 711 (54.4%): matched targeted therapy; 596 (45.6%) non-matched therapy.
- Median age: 57 yrs (range, 16-86); 39%, men.
- Median no. of prior therapies, 4 (range, 0-16); previously untreated = 2.8%
- Cancers: gastrointestinal, 24.2%; gynecological, 19.4%; breast, 13.5%; melanoma, 11.9%; lung, 8.7%.

| Response, evaluable               | Matched,<br>N = 697 | Non-matched,<br>N= 571 | <i>P</i> |
|-----------------------------------|---------------------|------------------------|----------|
| Objective response, %             | 16.2                | 5.4                    |          |
| Stable disease $\geq 6$ months, % | 18.7                | 14.7                   |          |
| Total, %                          | 34.9                | 20.1                   | <.001    |

# Progression-Free Survival by Type of Therapy



# Overall Survival by Type of Therapy



3-yr OS, 15% matched vs. 7% non-matched; 10-yr OS, 6% vs. 1%, respectively

# Precision Medicine in a Patient with Salivary Cancer (BRAF V600E Mutation, Vemurafenib)



## ***Taking aim sooner***

*If personalized medicine is to achieve its full potential, it should be used earlier on in clinical trials*

Many scientists ... believe that matching volunteers' genetic profiles to the drugs being tested will not only be better for the volunteers, but may also speed up the trials, and save millions of dollars in the process.

One such is Apostolia-Maria Tsimberidou of the University of Texas's MD Anderson Cancer Center, in Houston. And her preliminary results, presented at a meeting of the American Society of Clinical Oncology in Chicago, suggest she is right.



# Randomized Study Evaluating Molecular Profiling and Targeted Agents in Metastatic Cancer (IMPACT 2)

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## Primary Objective

To determine whether patients treated with a targeted therapy selected on the basis of mutational analysis of the tumor have longer **progression-free survival** from the time of randomization than those whose treatment is not selected based on alteration analysis

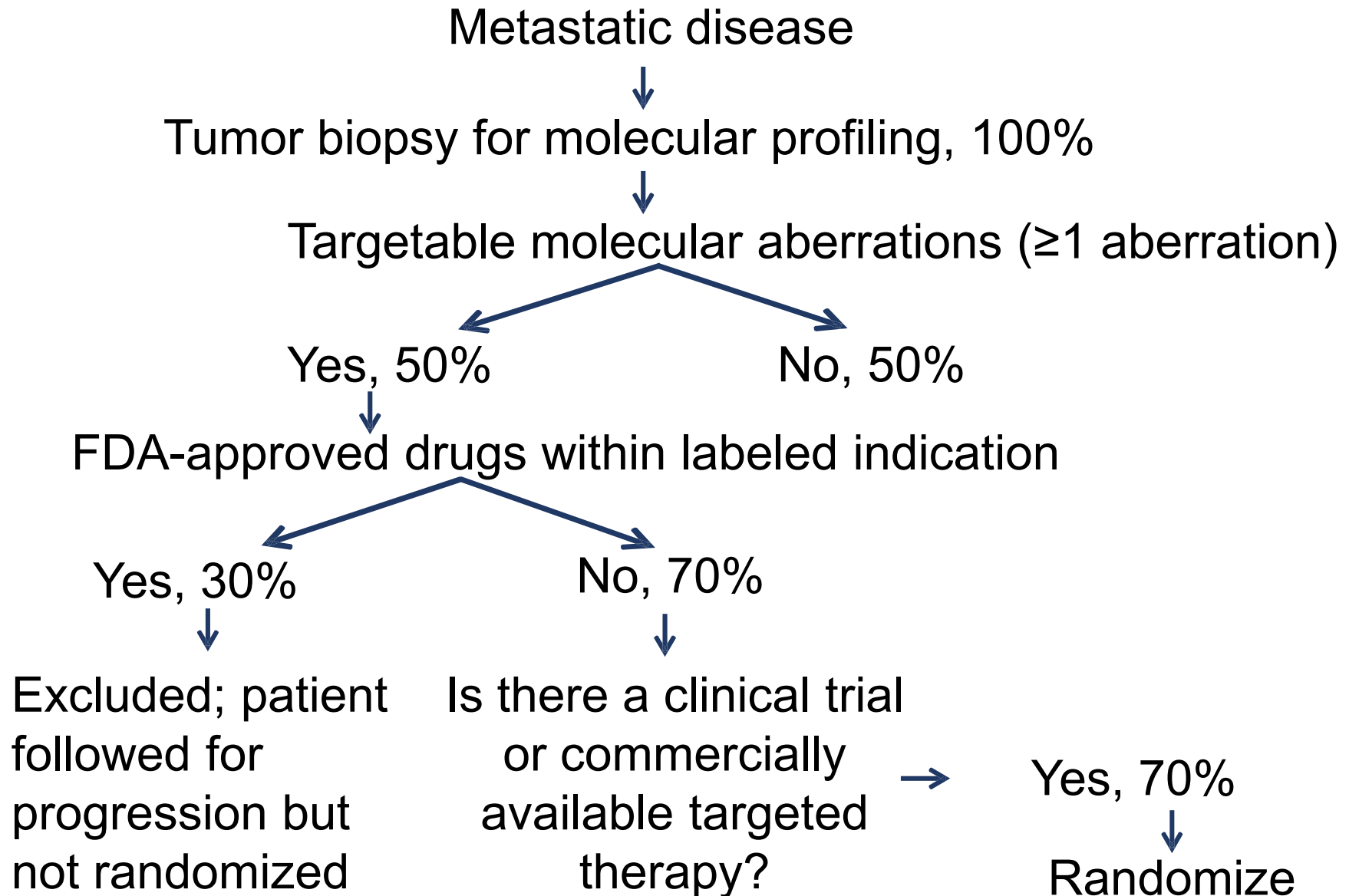
*PI: Tsimberidou, AM*

*[www.clinicaltrials.gov](https://www.clinicaltrials.gov) NCT02152254*

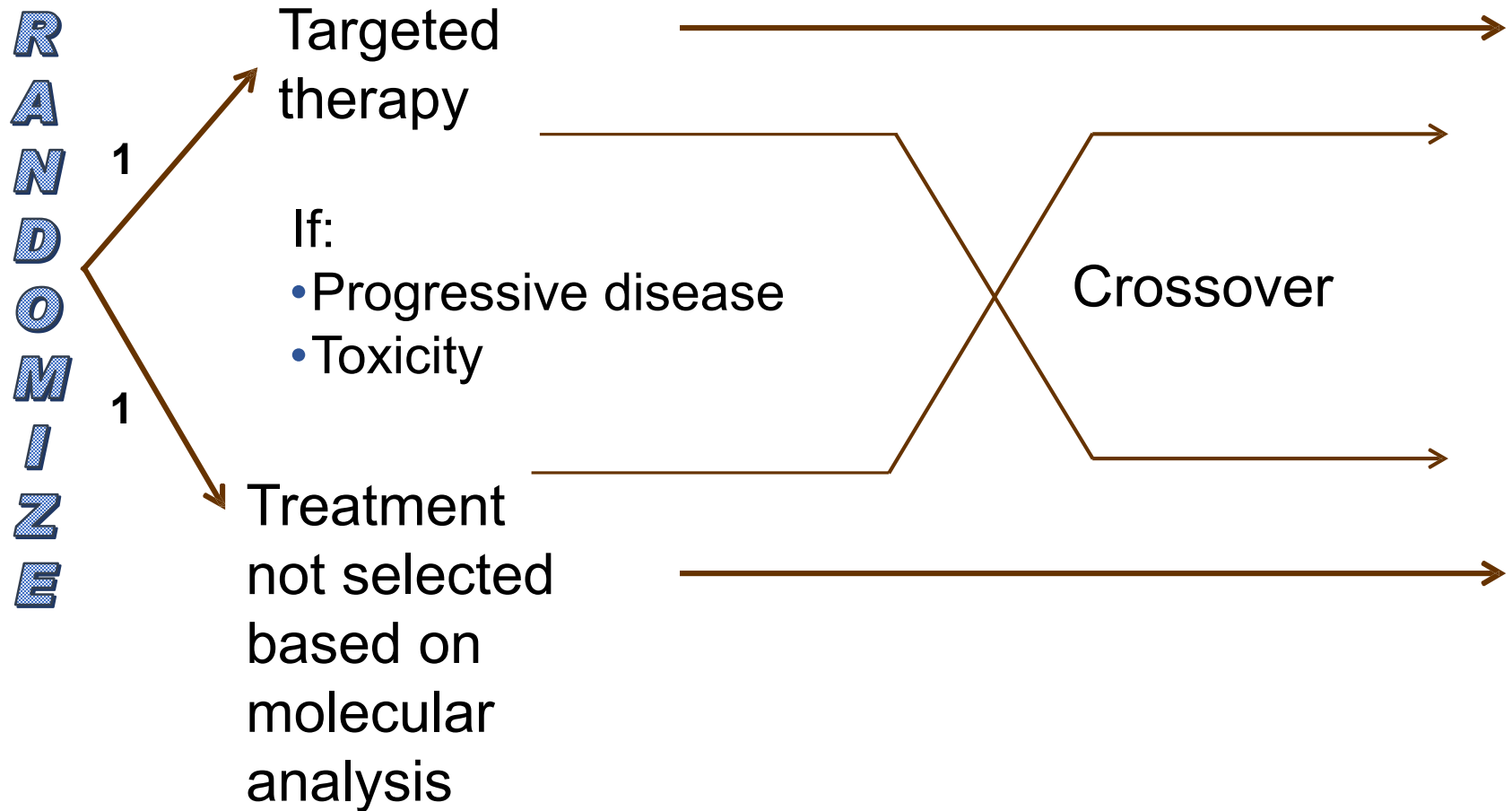
*Supported in part by a research grant, initially from Foundation Medicine and currently from Tempus*

# IMPACT 2. Study Design (I)

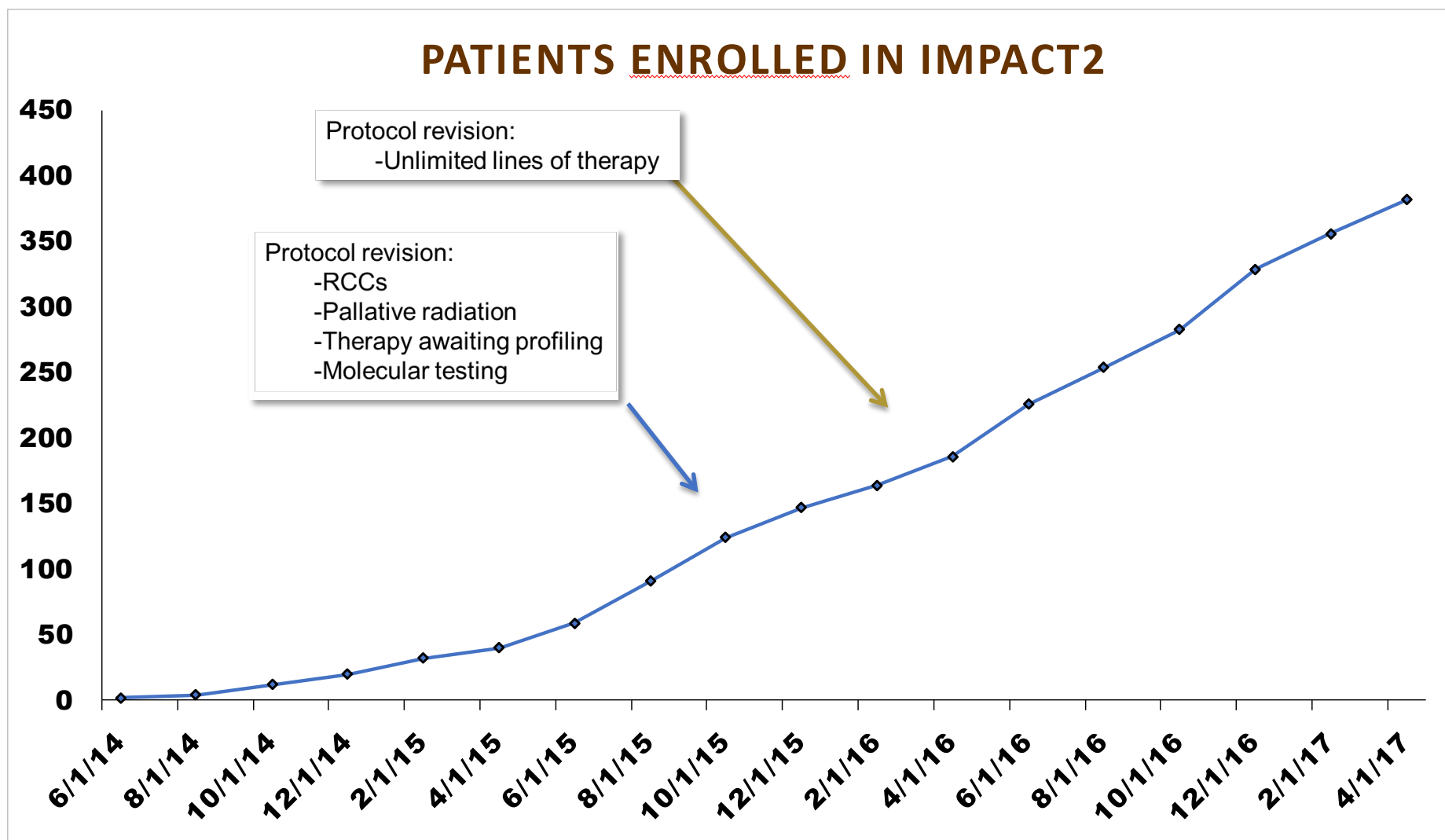
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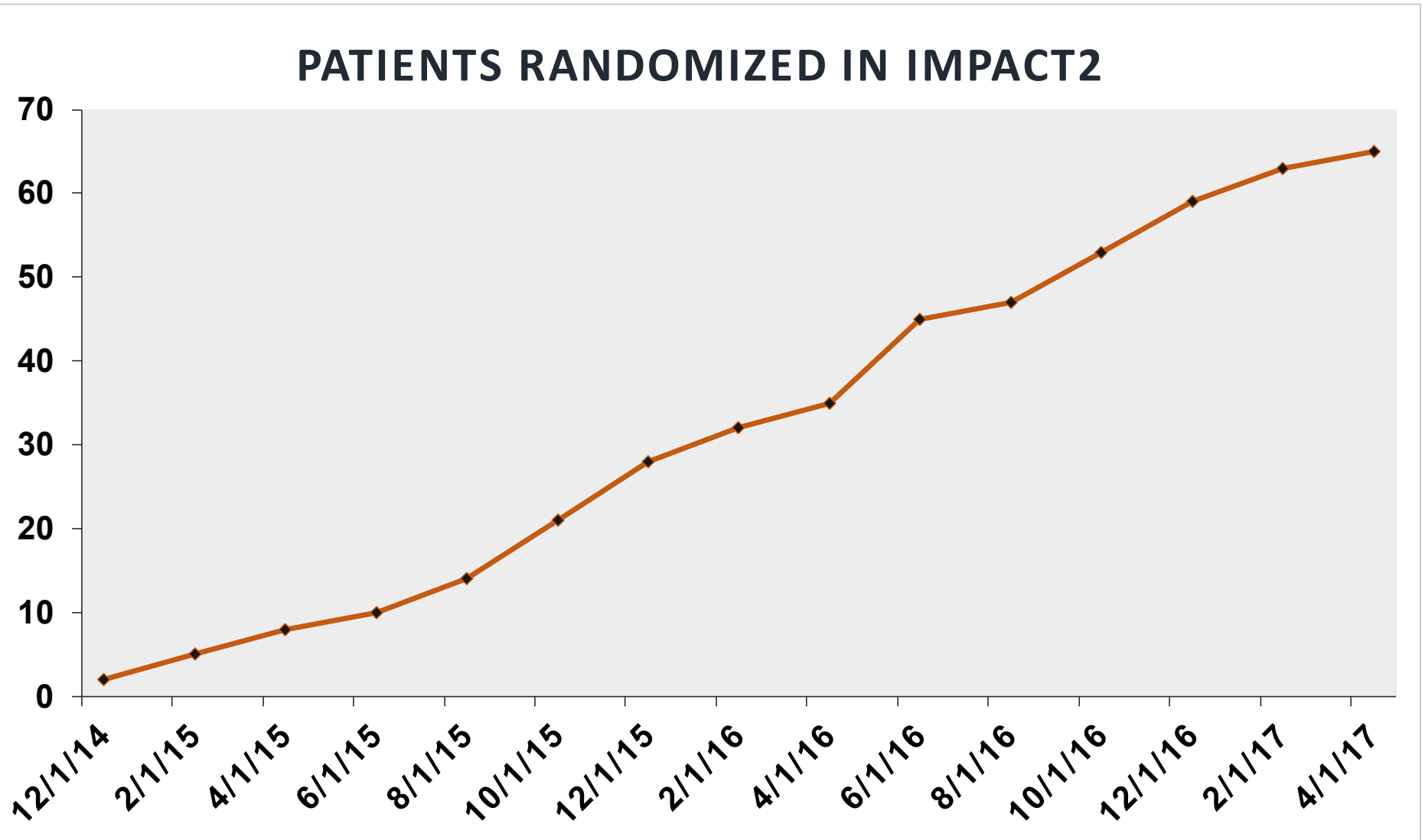
# IMPACT 2. Study Design (II)



# Cumulative plot of patients enrolled in IMPACT2



# Cumulative plot of patients randomized in IMPACT2



# Head and Neck Squamous Cell Carcinoma with *FGF* Amplifications: CR to FGFR Inhibitor

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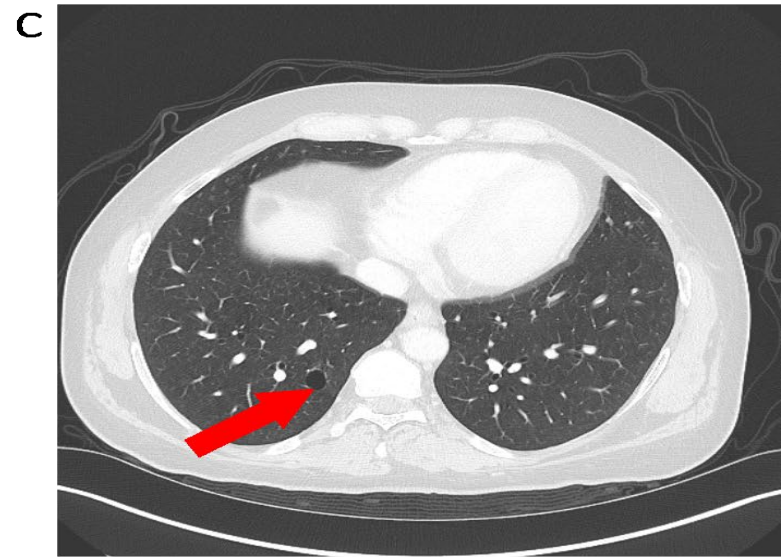
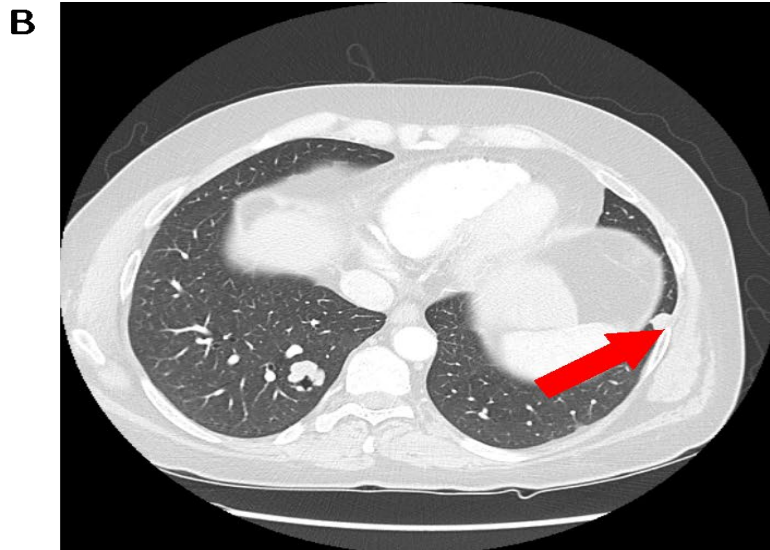
## Genomic Alterations

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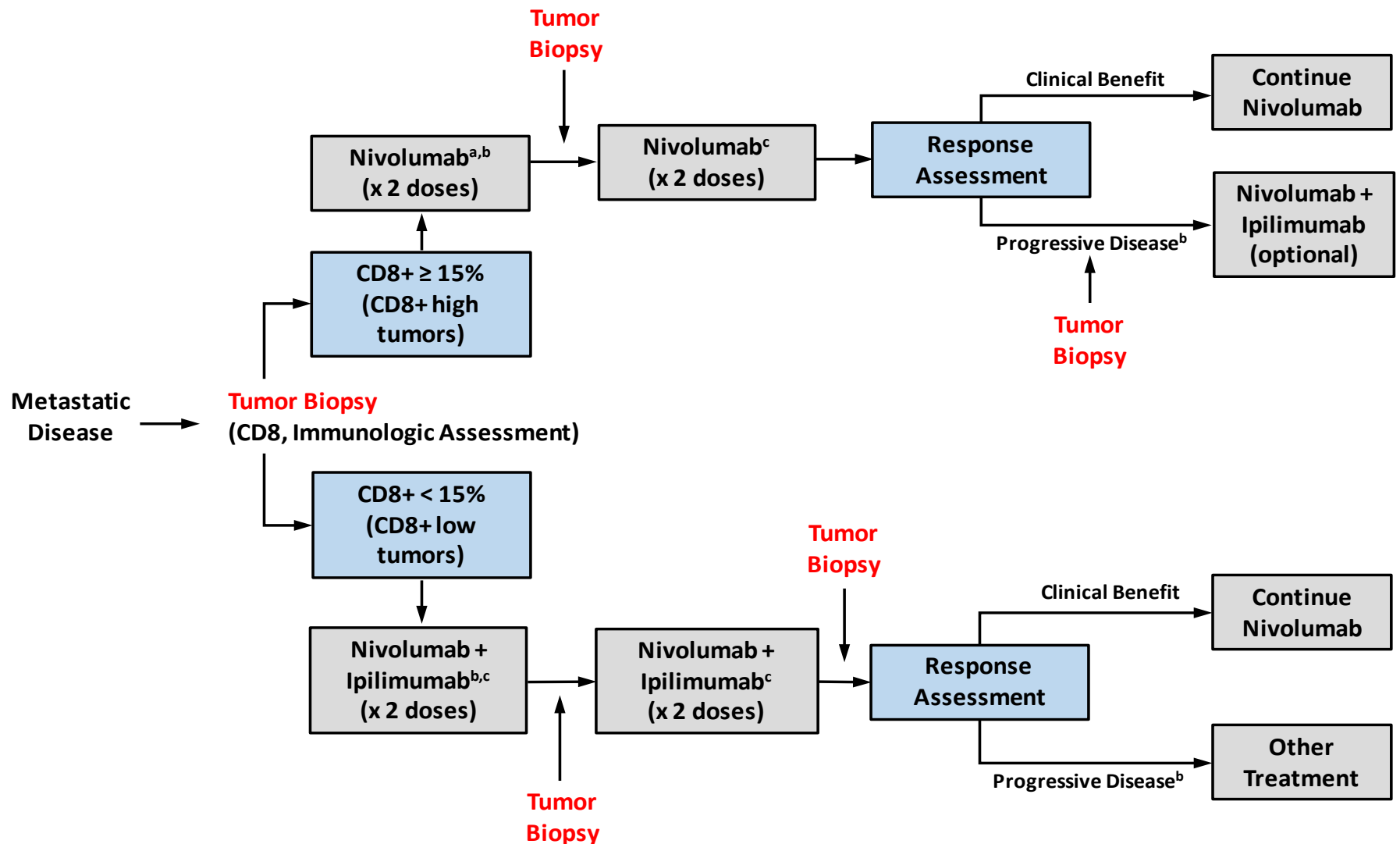
|                     |                           |
|---------------------|---------------------------|
| FGF19 amplification | CHD2 D213N                |
| FGF4 amplification  | CREBBP R1392*             |
| FGF23 amplification | EMSY amplification        |
| FGF3 amplification  | KDM5A amplification       |
| FGF6 amplification  | KRAS amplification        |
| CCND1 amplification | MYC duplication exons 2-3 |
| CCND2 amplification | TP53 E204*                |
| CDKN2A/B loss       |                           |

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# Head and Neck Squamous Cell Carcinoma with *FGF* Amplifications: CR to FGFR Inhibitor

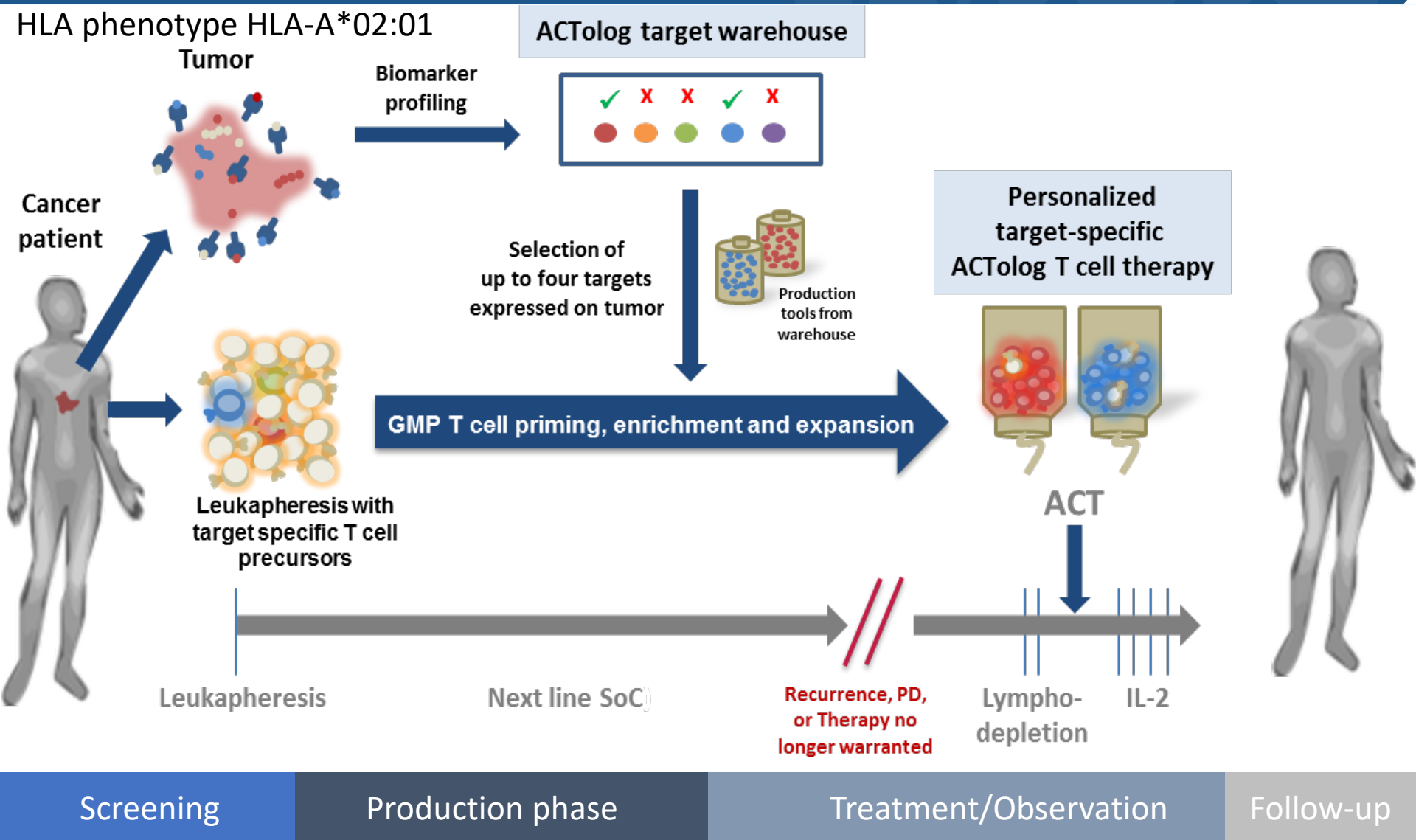


# An Exploratory Study of Nivolumab with or without Ipilimumab According to Tumor CD8 Expression in Patients with Advanced Cancer





# ACTolog: Endogenous CD8+ T cells in Advanced Cancer



# Challenges and Opportunities: Molecular profiling

|                         | Actual                     | Goal   |
|-------------------------|----------------------------|--|
| Tumor biopsy            | Not standard               | Standard of care   |
| Tumor sequencing        | Targeted NGS               | Whole genome sequencing, immune markers, transcriptomics, proteomics |
| Bioinformatics          | Limited                    | Optimized  |
| Emergence of sub-clones | Limited data               | Real-time monitoring   |
| Time to analysis        | >10 days                   | 1-3 days   |
| Timing                  | Advanced, refractory       | Starting at diagnosis  |
| Biomarker development   | Drug-specific              | Platform diagnostics   |
| Tumor heterogeneity     | Single lesion biopsy/ctDNA | Validated ctDNA analysis   |

# Challenges and Opportunities: Clinical Trials/Drugs

|                             | <b>Actual</b>  | <b>Goal</b>  |
|-----------------------------|--|--|
| Drug discovery              | Limited  | More, effective drugs                                |
| Study design                | Phase I, II, III   | Adaptive, “N of 1”, umbrella protocols               |
| Patient eligibility         | ≈5-30% of patients   | 100% of patients                                     |
| Histology-agnostic trial    | Small sample;<br>unbalanced data;<br>response<br>heterogeneity | Novel design for interim analyses; Adaptive design*  |
| “Targeted” drug definition  | Imprecise  | Precise  |
| Targeted therapy selection  | Subjective   | Evidence-based, tumor board, artificial intelligence |
| Adaptive learning, “N of 1” | <10%   | 100%   |

\* Early assessment of safety/clinical benefit of a drug permits inclusion of multiple stages of drug development in a single trial

# Faster, Better, Smarter Trials Require (I):

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- More patients to have “state of the art” comprehensive profiling (genomics, transcriptomics, immune markers, proteomics, novel markers)
- Profiling starting at diagnosis. Most studies offer drugs to patients who have received multiple lines of therapy; not a setting for optimal results
- More targeted, effective drugs/therapeutic strategies
- Information regarding available trials to doctors or patients to increase patient referral

# Faster, Better, Smarter Trials Require (II):

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- Better methodology: (current model: single drug for single aberration). Comprehensive profiling, clinical outcomes, advanced analyses:
  - (1) to offer matched therapy to more patients;
  - (2) to compare outcomes to those of patients not receiving matched therapy (case-control comparator) and
  - (3) to address complex questions that integrate precision molecular findings with immunologic findings to offer better treatments

# Implementation of Precision Medicine

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- Precision Medicine uses targeted therapy, immunotherapy, and other strategies to target specific biological abnormalities causing carcinogenesis in individual patients.
- Precision Cancer Medicine requires:
  1. Complete understanding of tumor biology, including immune features, that drives carcinogenesis
  2. Use of effective drugs and therapeutic strategies that inhibit carcinogenesis (rigorous definition)
  3. Access to testing and effective drugs for all patients starting at diagnosis and during the course of their disease

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**Thank you!**



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